

ABSTRACT

It is intended to provide a method whereby a protein can be separated and purified at a high accuracy by a convenient procedure. A
5 sample containing the desired protein is brought into contact with an ion exchanger under first conditions at a high ionic strength and at pH value not in the vicinity of the isoelectric point of the desired protein. Next, the component adsorbed by the ion exchanger is eluted under second conditions at a lower ionic strength than in the first
10 conditions and at a pH value closer to the isoelectric point of the desired protein than in the first conditions.